

## Problem Set

**Deadline:** Monday, Oct. 31, at 11:59pm.

**Submission:** You need to submit two files through Quercus:

- Your answers to the questions as a PDF file `problem_set_sol.pdf` (We encourage using  $\text{\LaTeX}$ ).
- Your implementation of question 3 as a Python file `code.py`.

Make sure your solutions and code structure are readable. You may lose up to 10 pts if we have a hard time reading your write-up/code.

## 1 Structure identifiability [20 pts]

Discovering causal graphs from observational distributions is generally impossible; we can identify the causal graph up to the skeleton and v-structures (see the observational equivalent theorem in lecture 2). For example, in the case of two dependent variables  $X \not\perp\!\!\!\perp Y$ , the causal graphs in Figure 1 are all feasible.

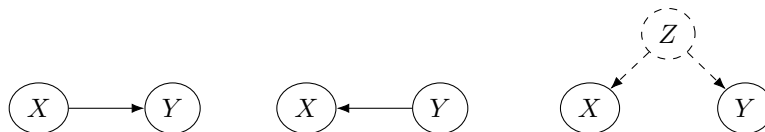


Figure 1

One approach to distinguish cause and effect is to make parametric assumptions on  $P_{\text{effect}|\text{cause}}$  and  $P_{\text{cause}}$ . For some of those parametric classes, there are theoretical results showing that there is only one direction ( $P_{Y|X}$  or  $P_{X|Y}$ ) that can satisfy the assumption. In this question, we will learn one such result for **linear** SCMs. Consider the following SCM, which induces joint distribution  $P(X, Y)$ :

$$Y := \alpha X + \epsilon_Y \quad X \perp\!\!\!\perp \epsilon_Y \quad (1.1)$$

for continuous 1-dimensional random variables  $Y, X, \epsilon_Y$ .

1. **[5 pts]** Assume  $X$  and  $\epsilon_Y$  are Gaussian. Then, find  $\beta \in \mathbb{R}$  such that the following SCM induces the same distribution  $P(X, Y)$ :

$$X := \beta Y + \epsilon_X \quad Y \perp\!\!\!\perp \epsilon_X \quad (1.2)$$

2. **[10 pts]** Now, assume there exists  $\beta \in \mathbb{R}$  and random variable  $\epsilon_X$  such that

$$X := \beta Y + \epsilon_X \quad Y \perp\!\!\!\perp \epsilon_X \quad (1.3)$$

generates the same distribution  $P(X, Y)$ . Show that  $X$  and  $\epsilon_Y$  are Gaussian.

*Hint: You can use the following theorem*

**Theorem 1.** Let  $X_1, X_2, \dots, X_n$  be  $n$  independent random variables, let

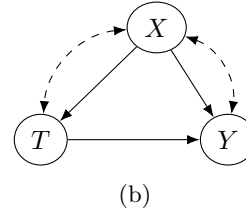
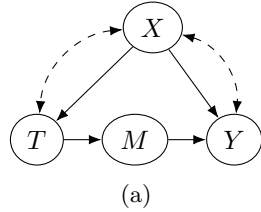
$$\begin{cases} Y_1 = \sum_{i=1}^n a_i X_i \\ Y_2 = \sum_{i=1}^n b_i X_i \end{cases} \quad (1.4)$$

and suppose  $Y_1$  and  $Y_2$  are independent. Now, if  $a_i b_i \neq 0$ , then  $X_i$  must be Gaussian.

3. [5 pts] Suppose we "believe" that the effect is a linear function of cause with non-Gaussian noise. (Informally) describe an algorithm that identifies cause and effect for two variables  $X$  and  $Y$ .

## 2 do-calculus! [20 pts]

Consider the following causal graphs.



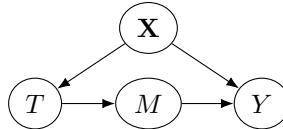
For each graph, is the causal effect of  $T$  on  $Y$  identifiable? If yes, use *do*-calculus and find a nonparametric formula based on observational distribution. If no, try to prove non-identifiability (e.g., using a counterexample).

## 3 Parametric estimation of causal effects [40 pts]

For this question, we have provided you with a simulated dataset, `data.csv`, of 5,000 samples with features  $\{Y, T, X_0, \dots, X_9\}$ . The goal is to estimate the average treatment effect on  $Y$ , i.e.,

$$\text{ATE} = \mathbb{E}[Y | do(T = 1)] - \mathbb{E}[Y | do(T = 0)] \quad (3.1)$$

Assume the corresponding causal graph as the following ( $\mathbf{X} = (X_0, \dots, X_9)$ ):

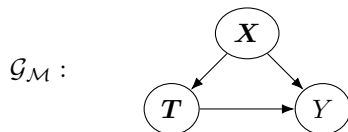


- Since the causal graph satisfies both Front-door and Back-door criteria, try to estimate the ATE using both formulas. In particular, you need to write three Python functions: `frontdoor`, `backdoor_com` (parametric Back-door formula with conditional outcome modeling), and `backdoor_grouped_com` (parametric Back-door formula with grouped conditional outcome modeling).
- Include the result of each of the three methods in your write-up. Also, report one estimation as your final answer. You can choose one of the three outputs, a combination of them, or just a random guess! (or any other causal estimation method). Part of your grading will be based on how close your final answer is to the true ATE.

- Make sure to include your implementation in a file `code.py`. Note that you can use any machine learning library to implement parametric functions for estimating conditional expectations, including linear regression, random forests, neural networks, etc. However, you *cannot* use libraries for causal estimation.

## 4 Can we identify unobserved confoundings? [20 pts]

Consider an SCM  $\mathcal{M} = (\mathbf{V}, \mathbf{U}, \mathcal{F}, P_{\mathbf{U}})$  with endogenous variables  $\mathbf{V} = \{X, \mathbf{T}, Y\}$  and unknown  $\mathcal{F}, P_{\mathbf{U}}$ . Assume the outcome and covariates are one-dimensional  $X, Y \in \mathbb{R}$ , and the treatment  $\mathbf{T} \in \mathbb{R}^m$ . In this question, we will focus on the Back-door causal graph, i.e.,



We know that in the case of unobserved confoundings  $X$ , the average treatment effect

$$\text{ATE} = \mathbb{E}[Y|do(\mathbf{T} = \mathbf{t})] - \mathbb{E}[Y|do(\mathbf{T} = \mathbf{t}')] \quad (4.1)$$

is not identifiable. In lecture 3, we showed that by constructing one example with  $m = 1$ . The reason for non-identifiability is that multiple choices of  $P(Y|X)$  and  $P(\mathbf{T}|X)$  can result in the same observed distribution  $P(\mathbf{T}, Y)$ . But, since we do not observe  $X$ , we cannot tell which one is the real one.

In this question, we want to test a hypothesis: *What if we can identify the latent factor?* In particular, we will consider the case of multi-dimensional treatments with  $m > 1$ . The intuition is that if we observe more causes, where each of them has some information about the unobserved confounding, it will be more likely to identify  $X$ . After identifying the confounder, we can substitute it with an estimation of it and use methods such as parametric G-formula to estimate the causal effect.

1. [5 pts] Let's formalize the above idea. To identify the hidden confounder, we can fit a probabilistic factor model<sup>1</sup> to capture the joint distribution of  $P(T_1, \dots, T_m)$ , where  $T_i$  is the  $i^{\text{th}}$  dimension of  $\mathbf{T}$ :

$$\begin{aligned} Z &\sim P_{\theta}(\cdot) \\ T_i|Z &\sim p_{\phi}(\cdot|Z) \quad i = 1, \dots, m, \end{aligned} \quad (4.2)$$

where  $Z$  is the latent variable. Suppose that, given  $m \gg 1$ , we can learn  $\theta$  and  $\phi$  such that the joint distribution of  $T_i$ s from eq. (4.2) matches the observed distribution of causes  $P(T_1, \dots, T_m)$ . Since  $T_i$ s are conditionally independent given  $Z$ , we can show their independence structure with the graphical model in Figure 3. Note that this graph is only to reason about conditional dependencies and not a causal graph.

Use  $d$ -separation to show that there cannot be a confounder  $X$  (other than  $Z$ ) that is the parent of multiple  $T_i$ s in Figure 3.

2. [5 pts] Given the result of the previous question, we can conclude that  $Z$  captures all the unobserved confounding. Since we can infer  $Z$ , the problem becomes a causal estimation with *observed* confounding. Describe an informal algorithm (using Back-door adjustment formula) to estimate the ATE using  $Z$ .

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<sup>1</sup>Some examples for factor models include [PCA](#), [VAEs](#), [ICA](#), [LDA](#), etc.

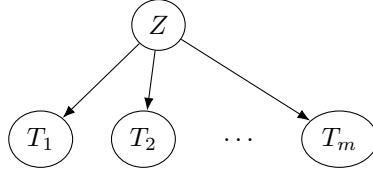


Figure 3

3. **[10 pts]** Show that the previous algorithm is wrong (!) by constructing a counterexample. In particular, show that the factorization of  $\mathbf{T}$  is not unique. You can construct two SCMs  $\mathcal{M}_1, \mathcal{M}_2$  ( $X, Y \in \mathbb{R}$  and  $\mathbf{T} \in \mathbb{R}^m$ ) with the same observed distribution  $P(\mathbf{T}, Y)$  and different ATEs, where the treatments can be factorized in both, i.e.,

$$\begin{aligned}
 P_{\mathcal{M}_1}(T_1, \dots, T_m | X) &= \prod_{i=1}^m P_{\mathcal{M}_1}(T_i | X) \\
 P_{\mathcal{M}_2}(T_1, \dots, T_m | X) &= \prod_{i=1}^m P_{\mathcal{M}_2}(T_i | X)
 \end{aligned} \tag{4.3}$$

If you cannot solve it for the general case, try constructing a numerical example for  $m = 3$ .

*Hint: You can consider linear SCMs with Gaussian noise in the form of*

$$\begin{aligned}
 X &:= \epsilon_X \\
 \mathcal{M}: \quad \mathbf{T} &:= \alpha X + \boldsymbol{\epsilon}_{\mathbf{T}} \\
 Y &:= \boldsymbol{\beta}^\top \mathbf{T} + \gamma X + \epsilon_Y
 \end{aligned} \tag{4.4}$$

with  $\epsilon_X \sim \mathcal{N}(0, \sigma_X)$ ,  $\epsilon_Y \sim \mathcal{N}(0, \sigma_Y)$ , and  $\boldsymbol{\epsilon}_{\mathbf{T}} \sim \mathcal{N}(\mathbf{0}, \boldsymbol{\Sigma}_{\mathbf{T}})$  ( $\boldsymbol{\Sigma}_{\mathbf{T}}$  is a diagonal  $m \times m$  covariance matrix). First, you need to show that  $X$  factorizes  $P(\mathbf{T})$  in this SCM. Then, construct a new SCM with different coefficients and the same induced distribution  $P(\mathbf{T}, Y)$ . You can try changing the treatment assignment to

$$\mathbf{T} := c \cdot \alpha X + \boldsymbol{\epsilon}_{\mathbf{T}} \tag{4.5}$$

and see how other coefficients should be changed.